

50C

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/516,490A
Source: PG 1/0
Date Processed by STIC: 1/10/06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.2.2 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05



PCT

RAW SEQUENCE LISTING

DATE: 01/10/2006

PATENT APPLICATION: US/10/516,490A

TIME: 08:39:08

Input Set : A:\X15642.NatlPhase.ST25.txt

Output Set: N:\CRF4\01102006\J516490A.raw

3 <110> APPLICANT: Richard Dennis DiMarchi
 4 David Lee Smiley
 5 Lianshan Zhang
 7 <120> TITLE OF INVENTION: MODIFIED GLUCAGON-LIKE PEPTIDE-1 ANALOGS
 9 <130> FILE REFERENCE: X-15642 National Phase
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/516,490A
 C--> 11 <141> CURRENT FILING DATE: 2004-12-01
 11 <160> NUMBER OF SEQ ID NOS: 24
 13 <170> SOFTWARE: PatentIn version 3.2
 15 <210> SEQ ID NO: 1
 16 <211> LENGTH: 31
 17 <212> TYPE: PRT
 18 <213> ORGANISM: Artificial
 20 <220> FEATURE:
 21 <223> OTHER INFORMATION: Synthetic constructs
 24 <220> FEATURE:
 25 <221> NAME/KEY: MISC_FEATURE
 26 <222> LOCATION: (1)..(1) ✓
 27 <223> OTHER INFORMATION: Xaa= L-histidine, D-histidine, desamino-histidine,
 28 2-amino-histidine, beta-hydroxy-
 29 histidine, homohistidine, alpha-fluoromethyl-histidine, or alpha
 30 methyl-histidine
 32 <220> FEATURE:
 33 <221> NAME/KEY: MISC_FEATURE
 34 <222> LOCATION: (2)..(2) ✓
 35 <223> OTHER INFORMATION: Xaa= Ala, Gly, Val, Leu, Ile, Ser, or Thr
 37 <220> FEATURE:
 38 <221> NAME/KEY: MISC_FEATURE
 39 <222> LOCATION: (6)..(6) ✓
 40 <223> OTHER INFORMATION: Xaa= Phe, Trp, or Tyr
 42 <220> FEATURE:
 43 <221> NAME/KEY: MISC_FEATURE
 44 <222> LOCATION: (10)..(10) ✓
 45 <223> OTHER INFORMATION: Xaa= Val, Trp, Ile, Leu, Phe, or Tyr
 47 <220> FEATURE:
 48 <221> NAME/KEY: MISC_FEATURE
 49 <222> LOCATION: (12)..(12) ✓
 50 <223> OTHER INFORMATION: Xaa= Ser, Trp, Tyr, Phe, Lys, Ile, Leu, Val
 52 <220> FEATURE:
 53 <221> NAME/KEY: MISC_FEATURE
 54 <222> LOCATION: (13)..(13) ✓
 55 <223> OTHER INFORMATION: Xaa= Tyr, Trp, or Phe
 57 <220> FEATURE:

2-3, 5-7
**Does Not Comply
 Corrected Diskette Needed**

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/516,490A

DATE: 01/10/2006

TIME: 08:39:08

Input Set : A:\X15642.Nat1Phase.ST25.txt
 Output Set: N:\CRF4\01102006\J516490A.raw

```

58 <221> NAME/KEY: MISC_FEATURE
59 <222> LOCATION: (14)..(14)
60 <223> OTHER INFORMATION: Xaa= Leu, Phe, Tyr, or Trp
62 <220> FEATURE:
63 <221> NAME/KEY: MISC_FEATURE
64 <222> LOCATION: (16)..(16)
65 <223> OTHER INFORMATION: Xaa= Gly, Glu, Asp, Lys
67 <220> FEATURE:
68 <221> NAME/KEY: MISC_FEATURE
69 <222> LOCATION: (19)..(19)
70 <223> OTHER INFORMATION: Xaa= Ala, Val, Ile, or Leu
72 <220> FEATURE:
73 <221> NAME/KEY: MISC_FEATURE
74 <222> LOCATION: (21)..(21)
75 <223> OTHER INFORMATION: Xaa= Glu, Ile, or Ala
77 <220> FEATURE:
78 <221> NAME/KEY: MISC_FEATURE
79 <222> LOCATION: (24)..(24)
80 <223> OTHER INFORMATION: Xaa= Ala, or Glu
82 <220> FEATURE:
83 <221> NAME/KEY: MISC_FEATURE
84 <222> LOCATION: (27)..(27)
85 <223> OTHER INFORMATION: Xaa= Val, or Ile
87 <220> FEATURE:
88 <221> NAME/KEY: MISC_FEATURE
89 <222> LOCATION: (31)..(31)
90 <223> OTHER INFORMATION: Xaa= L-Cys, D-Cys, homocysteine, or penicillamine
92 <400> SEQUENCE: 1
W--> 94 Xaa Xaa Glu Gly Thr Xaa Thr Ser Asp Xaa Ser Xaa Xaa Xaa Glu Xaa
      95 1          5          10          15
W--> 98 Gln Ala Xaa Lys Xaa Phe Ile Xaa Trp Leu Xaa Lys Gly Arg Xaa
      99          20          25          30
102 <210> SEQ ID NO: 2
103 <211> LENGTH: 31
104 <212> TYPE: PRT
105 <213> ORGANISM: Artificial
107 <220> FEATURE:
108 <223> OTHER INFORMATION: Synthetic construct
111 <220> FEATURE:
112 <221> NAME/KEY: MISC_FEATURE
113 <222> LOCATION: (1)..(1)
114 <223> OTHER INFORMATION: Xaa= L-histidine, D-histidine, desamino-
      histidine,
115      2-amino-histidine, beta-hydroxy-
116      histidine, homohistidine, alpha-fluoromethyl-histidine, or
117      alpha-methyl-histidine
119 <220> FEATURE:
120 <221> NAME/KEY: MISC_FEATURE
121 <222> LOCATION: (2)..(2)
122 <223> OTHER INFORMATION: Xaa= Gly, Ala, Val, Leu, Ile, Ser or Thr

```

RAW SEQUENCE LISTING

DATE: 01/10/2006

PATENT APPLICATION: US/10/516,490A

TIME: 08:39:08

Input Set : A:\X15642.Nat1Phase.ST25.txt

Output Set: N:\CRF4\01102006\J516490A.raw

124 <220> FEATURE:
 125 <221> NAME/KEY: MISC_FEATURE
 126 <222> LOCATION: (10)..(10)
 127 <223> OTHER INFORMATION: Xaa = Val, Phe, Tyr, or Trp
 129 <220> FEATURE:
 130 <221> NAME/KEY: MISC_FEATURE
 131 <222> LOCATION: (12)..(12)
 132 <223> OTHER INFORMATION: Xaa = Ser, Tyr, Trp, Phe, Lys, Ile, Leu, or Val
 134 <220> FEATURE:
 135 <221> NAME/KEY: MISC_FEATURE
 136 <222> LOCATION: (16)..(16)
 137 <223> OTHER INFORMATION: Xaa = Gly, Clu Asp, or Lys *do you mean Glu?*
 139 <220> FEATURE:
 140 <221> NAME/KEY: MISC_FEATURE
 141 <222> LOCATION: (19)..(19)
 142 <223> OTHER INFORMATION: Xaa = Ala, Val, Ile, or Leu
 144 <220> FEATURE:
 145 <221> NAME/KEY: MISC_FEATURE
 146 <222> LOCATION: (27)..(27)
 147 <223> OTHER INFORMATION: Xaa = Val or Ile
 149 <220> FEATURE:
 150 <221> NAME/KEY: MISC_FEATURE
 151 <222> LOCATION: (31)..(31)
 152 <223> OTHER INFORMATION: Xaa = L-Cys, D-Cys, homocysteine, or
 penicillamine
 154 <400> SEQUENCE: 2
 W--> 156 Xaa Xaa Glu Gly Thr Phe Thr Ser Asp Xaa Ser Xaa Tyr Leu Glu Xaa
 157 1 5 10 15
 W--> 160 Gln Ala Xaa Lys Glu Phe Ile Ala Trp Leu Xaa Lys Gly Arg Xaa
 161 20 25 30
 164 <210> SEQ ID NO: 3
 165 <211> LENGTH: 42
 166 <212> TYPE: PRT
 167 <213> ORGANISM: Artificial
 169 <220> FEATURE:
 170 <223> OTHER INFORMATION: Synthetic construct
 173 <220> FEATURE:
 174 <221> NAME/KEY: MISC_FEATURE
 175 <222> LOCATION: (1)..(1)
 176 <223> OTHER INFORMATION: Xaa = L-histidine, D-histidine, desamino-
 histidine,
 177 2-amino-histidine, beta-hydroxy-
 178 histidine, homohistidine, alpha-fluoromethyl-histidine, or
 179 alpha-methyl-histidine
 181 <220> FEATURE:
 182 <221> NAME/KEY: MISC_FEATURE
 183 <222> LOCATION: (2)..(2)
 184 <223> OTHER INFORMATION: Xaa = Ala, Gly, Val, Leu, Ile, Ser, or Thr
 186 <220> FEATURE:
 187 <221> NAME/KEY: MISC_FEATURE
 188 <222> LOCATION: (6)..(6)

RAW SEQUENCE LISTING

DATE: 01/10/2006

PATENT APPLICATION: US/10/516,490A

TIME: 08:39:08

Input Set : A:\X15642.Nat1Phase.ST25.txt

Output Set: N:\CRF4\01102006\J516490A.raw

```

189 <223> OTHER INFORMATION: Xaa = Phe, Trp, or Tyr
191 <220> FEATURE:
192 <221> NAME/KEY: MISC_FEATURE
193 <222> LOCATION: (10)..(10)
194 <223> OTHER INFORMATION: Xaa = Val, Trp, Ile, Leu, Phe, or Tyr
196 <220> FEATURE:
197 <221> NAME/KEY: MISC_FEATURE
198 <222> LOCATION: (12)..(12)
199 <223> OTHER INFORMATION: Xaa = Ser, Trp, Tyr, Phe, Lys, Ile, Leu, Val
201 <220> FEATURE:
202 <221> NAME/KEY: MISC_FEATURE
203 <222> LOCATION: (13)..(13)
204 <223> OTHER INFORMATION: Xaa = Tyr, Trp, or Phe
206 <220> FEATURE:
207 <221> NAME/KEY: MISC_FEATURE
208 <222> LOCATION: (14)..(14)
209 <223> OTHER INFORMATION: Xaa = Leu, Phe, Tyr, or Trp
211 <220> FEATURE:
212 <221> NAME/KEY: MISC_FEATURE
213 <222> LOCATION: (16)..(16)
214 <223> OTHER INFORMATION: Xaa = Gly, Glu, Asp, or Lys
216 <220> FEATURE:
217 <221> NAME/KEY: MISC_FEATURE
218 <222> LOCATION: (19)..(19)
219 <223> OTHER INFORMATION: Xaa = Ala, Val, Ile, or Leu
221 <220> FEATURE:
222 <221> NAME/KEY: MISC_FEATURE
223 <222> LOCATION: (21)..(21)
224 <223> OTHER INFORMATION: Xaa = Glu, Ile, or Ala
226 <220> FEATURE:
227 <221> NAME/KEY: MISC_FEATURE
228 <222> LOCATION: (24)..(24)
229 <223> OTHER INFORMATION: Xaa = Ala or Glu
231 <220> FEATURE:
232 <221> NAME/KEY: MISC_FEATURE
233 <222> LOCATION: (27)..(27)
234 <223> OTHER INFORMATION: Xaa = Val or Ile
236 <220> FEATURE:
237 <221> NAME/KEY: MISC_FEATURE
238 <222> LOCATION: (28)..(28)
239 <223> OTHER INFORMATION: Xaa = Lys, Asp, Arg, or Glu
241 <220> FEATURE:
242 <221> NAME/KEY: MISC_FEATURE
243 <222> LOCATION: (30)..(30)
244 <223> OTHER INFORMATION: Xaa = Gly, Pro, or Arg
246 <220> FEATURE:
247 <221> NAME/KEY: MISC_FEATURE
248 <222> LOCATION: (31)..(31)
249 <223> OTHER INFORMATION: Xaa = Gly, Pro, Ser, L-Cys, D-Cys, homocysteine,
or penicillamine

```

RAW SEQUENCE LISTING

DATE: 01/10/2006

PATENT APPLICATION: US/10/516,490A

TIME: 08:39:08

Input Set : A:\X15642.NatlPhase.ST25.txt

Output Set: N:\CRF4\01102006\J516490A.raw

251 <220> FEATURE:
 252 <221> NAME/KEY: MISC_FEATURE ✓
 253 <222> LOCATION: (32)..(32)
 254 <223> OTHER INFORMATION: Xaa = Ser, Pro, His, L-Cys, D-Cys, homocysteine,
 penicillamine, *do you mean "amidation"? If so, please add "amidated" to "NH₂"*
 255 NH₂
 257 <220> FEATURE:
 258 <221> NAME/KEY: MISC_FEATURE ✓
 259 <222> LOCATION: (33)..(33)
 260 <223> OTHER INFORMATION: Xaa = Ser, Arg, Thr, Trp, Lys, L-Cys, D-Cys,
 homocysteine,
 261 penicillamine, NH₂ or
 262 is absent
 264 <220> FEATURE:
 265 <221> NAME/KEY: MISC_FEATURE ✓
 266 <222> LOCATION: (34)..(34)
 267 <223> OTHER INFORMATION: Xaa = Ser, Gly, L-Cys, D-Cys, homocysteine,
 penicillamine, NH₂,
 268 or is absent
 270 <220> FEATURE:
 271 <221> NAME/KEY: MISC_FEATURE ✓
 272 <222> LOCATION: (35)..(35)
 273 <223> OTHER INFORMATION: Xaa = Ala, Asp, Arg, Glu, Lys, Gly, L-Cys, D-Cys,
 homocysteine,
 274 penicillamine,
 275 NH₂ or is absent
 277 <220> FEATURE:
 278 <221> NAME/KEY: MISC_FEATURE ✓
 279 <222> LOCATION: (36)..(36)
 280 <223> OTHER INFORMATION: Xaa = Pro, Ala, L-Cys, D-Cys, homocysteine,
 penicillamine, NH₂,
 281 or is absent
 283 <220> FEATURE:
 284 <221> NAME/KEY: MISC_FEATURE ✓
 285 <222> LOCATION: (37)..(37)
 286 <223> OTHER INFORMATION: Xaa = Pro, Ala, L-Cys, D-Cys, homocysteine,
 penicillamine, NH₂ or
 287 is absent
 289 <220> FEATURE:
 290 <221> NAME/KEY: MISC_FEATURE ✓
 291 <222> LOCATION: (38)..(38)
 292 <223> OTHER INFORMATION: Xaa = Pro, Ala, Arg, Lys, His, L-Cys, D-Cys, *Cys?*
 homocysteine,
 293 penicillamine, NH₂ or
 294 is absent
 296 <220> FEATURE:
 297 <221> NAME/KEY: MISC_FEATURE ✓
 298 <222> LOCATION: (39)..(39)
 299 <223> OTHER INFORMATION: Xaa = Ser, His, Pro, Lys, Arg, L-Cys, D-Cys,
 homocysteine,
 300 penicillamine, NH₂ or
 301 is absent
 303 <220> FEATURE:

304 <221> NAME/KEY: MISC_FEATURE
305 <222> LOCATION: (40)..(40) ✓
306 <223> OTHER INFORMATION: Xaa = His, Ser, Arg, Lys, L-Cys, D-Cys,
homocysteine,
307 penicillamine, NH₂ or

*Please ensure that amino acids are
spelled correctly in subsequent sequences.*

RAW SEQUENCE LISTING ERROR SUMMARY
 PATENT APPLICATION: US/10/516,490A

DATE: 01/10/2006
 TIME: 08:39:09

Input Set : A:\X15642.NatlPhase.ST25.txt
 Output Set: N:\CRF4\01102006\J516490A.raw

FYI

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220>

to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1,2,6,10,12,13,14,16,19,21,24,27,31
 Seq#:2; Xaa Pos. 1,2,10,12,16,19,27,31
 Seq#:3; Xaa Pos. 1,2,6,10,12,13,14,16,19,21,24,27,28,30,31,32,33,34,35,36
 Seq#:3; Xaa Pos. 37,38,39,40,41,42
 Seq#:4; Xaa Pos. 1,2,10,16,19,27,28,30,31,32,33,34,35,36,37,38,39,40,41,42
 Seq#:5; Xaa Pos. 1,2,16,19,27,32,33,34,35,36,37,38,39,40,41,42
 Seq#:6; Xaa Pos. 1,2,6,10,12,13,14,16,19,21,24,27,28,30,31,32,33,34,35,36
 Seq#:6; Xaa Pos. 37,38,39,40,41,42,43,44,45
 Seq#:7; Xaa Pos. 32,33,34,35,36,37,38,39,40,41,42,43,44,45
 Seq#:8; Xaa Pos. 1,2,6,10,12,13,14,16,19,21,24,27
 Seq#:9; Xaa Pos. 1,2,10,12,16,19,27
 Seq#:10; Xaa Pos. 1,2,6,10,12,13,14,16,19,21,24,27,28,30,31,32,33,34,35,36
 Seq#:10; Xaa Pos. 37,38,39,40,41,42
 Seq#:11; Xaa Pos. 1,2,10,16,19,27,28,30,31,32,33,34,35,36,37,38,39,40,41,42
 Seq#:12; Xaa Pos. 1,2,16,19,27,32,33,34,35,36,37,38,39,40,41,42
 Seq#:13; Xaa Pos. 1,2,6,10,12,13,14,16,19,21,24,27,28,30,31,32,33,34,35,36
 Seq#:13; Xaa Pos. 37,38,39,40,41,42,43,44,45
 Seq#:14; Xaa Pos. 32,33,34,35,36,37,38,39,40,41,42,43,44,45
 Seq#:15; Xaa Pos. 1,2,6,10,12,13,14,16,19,21,24,27,31

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22

VERIFICATION SUMMARY

DATE: 01/10/2006

PATENT APPLICATION: US/10/516,490A

TIME: 08:39:09

Input Set : A:\X15642.NatlPhase.ST25.txt

Output Set: N:\CRF4\01102006\J516490A.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:94 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:98 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:16
L:156 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:160 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:16
L:324 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:328 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:16
L:332 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:32
L:466 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:470 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:16
L:474 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:32
L:583 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:587 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:16
L:591 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:32
L:762 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:766 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:16
L:770 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:32
L:869 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:16
L:873 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:32
L:951 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:955 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:16
L:1008 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:1012 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:16
L:1160 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:1164 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:16
L:1168 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:32
L:1286 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:1290 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:16
L:1294 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:32
L:1391 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:1395 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:16
L:1399 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:32
L:1561 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
L:1565 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:16
L:1569 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:32
L:1658 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:16
L:1662 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:32
L:1745 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:1749 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:16